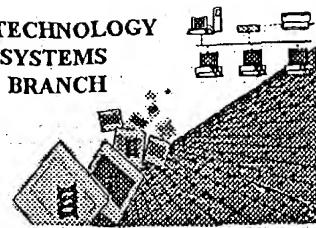# RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _09/462,480A_

Source: _1FW16_

Date Processed by STIC: _11/23/04_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.2 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
      http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission User Manual - ePAVE).
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04): U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04

Raw Sequence Listing Error Summary

ERROR DETECTED   **SUGGESTED CORRECTION**   SERIAL NUMBER: _09/462,480A_

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 ____ Wrapped Nucleics
Wrapped Aminos
The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 ____ Invalid Line Length
The rules require that a line **not exceed** 72 characters in length. This includes white spaces.

3 ____ Misaligned Amino
Numbering
The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.

4 ____ Non-ASCII
The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**

5 ____ Variable Length
Sequence(s) ____ contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 ____ PatentIn 2.0
"bug"
A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

7 ____ Skipped Sequences
(OLD RULES)
Sequence(s) ____ missing. If intentional, please insert the following lines for **each** skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8 ____ Skipped Sequences
(NEW RULES)
Sequence(s) ____ missing. **If intentional,** please insert the following lines for **each** skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 ____ Use of n's or Xaa's
(NEW RULES)
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 ____ Invalid <213>
Response
Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence *Do not Combine responses.*

11 ____ Use of <220>
Sequence(s) ____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32)   (Sec. 1.823 of Sequence Rules)

12 ____ PatentIn 2.0
"bug"
Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 ____ Misuse of n/Xaa
"n" can **only** represent a single <u>nucleotide</u>; "Xaa" can **only** represent a single <u>amino acid</u>

AMC – Biotechnology Systems Branch – 09/09/2003

IFW16

RAW SEQUENCE LISTING   DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A   TIME: 14:08:31

Input Set : A:\066001650XPCT.txt
Output Set: N:\CRF4\11222004\I462480A.raw

```
 3 <110> APPLICANT: GICQUEL, BRIGITTE
 4       BERTHET, FRANCIOS-XAVIER
 5       ANDERSEN, PETER
 6       RASMUSSEN, PETER BIRK
 8 <120> TITLE OF INVENTION: POLYNUCLEOTIDE FUNCTIONALLY CODING FOR THE LHP PROTEIN FROM
 9       MYCOBACTERIUM TUBERCULOSIS, ITS BIOLOGICALLY ACTIVE DERIVATIVE
10       FRAGMENTS, AS WELL AS METHODS USING THE SAME
12 <130> FILE REFERENCE: 0660-0165-0XPCT
14 <140> CURRENT APPLICATION NUMBER: 09/462,480A
15 <141> CURRENT FILING DATE: 2000-03-06
17 <150> PRIOR APPLICATION NUMBER: PCT/IB98/01091
18 <151> PRIOR FILING DATE: 1998-07-16
20 <150> PRIOR APPLICATION NUMBER: 60/052,631
21 <151> PRIOR FILING DATE: 1997-07-16
23 <160> NUMBER OF SEQ ID NOS: 34
25 <170> SOFTWARE: PatentIn version 3.3
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 1277
29 <212> TYPE: DNA
30 <213> ORGANISM: Mycobacterium tuberculosis
32 <400> SEQUENCE: 1
33 ctgcagcagg tgacgtcgtt gttcagccag gtgggcggca ccggcggcgg caacccagcc    60
35 gacgaggaag ccgcgcagat gggcctgctc ggcaccagtc cgctgtcgaa ccatccgctg   120
37 gctggtggat caggccccag cgcgggcgcg ggcctgctgc gcgcggagtc gctacctggc   180
39 gcaggtgggt cgttgacccg cacgccgctg atgtctcagc tgatcgaaaa gccggttgcc   240
41 ccctcggtga tgccggcggc tgttgccgga tcgtcggtga cgggtggcgc cgctccggtg   300
43 ggtccgggag cgatgggcca gggttcgcaa tccggcggct ccaccagccc gggtctggtc   360
45 gcgccggcac cgctcgcgca ggagcgtgaa gaagacgacg aggacgactg ggacgaagag   420
47 gacgactggt gagctcccgt aatgacaaca gacttcccgg ccacccgggc cggaagactt   480
49 gccaacattt tggcgaggaa ggtaaagaga gaaagtagtc cagcatggca gagatgaaga   540
51 ccgatgccgc taccctcggg caggaggcag gtaatttcga gcggatctcc ggcgacctga   600
53 aaacccagat cgaccaggtg gagtcgacgg caggttcgtt gcagggccag tggcgcggcg   660
55 cggcggggac ggccgcccag gccgcggtgg tgcgcttcca agaagcagcc aataagcaga   720
57 agcaggaact cgacgagatc tcgacgaata ttcgtcaggc cggcgtccaa tactcgaggg   780
59 ccgacgagga gcagcagcag gcgctgtcct cgcaaatggg cttctgaccc gctaatacga   840
61 aaagaaacgg agcaaaaaca tgacagagca gcagtggaat ttcgcgggta tcgaggccgc   900
63 ggcaagcgca atccagggaa atgtcacgtc cattcattcc ctccttgacg aggggaagca   960
65 gtccctgacc aagctcgcag cggcctgggg cggtagcggt tcgaggcgt accagggtgt  1020
67 ccagcaaaaa tgggacgcca cggctaccga gctgaacaac gcgctgcaga acctggcgcg  1080
69 gacgatcagc gaagccggtc aggcaatggc ttcgaccgaa ggcaacgtca ctgggatgtt  1140
71 cgcatagggc aacgccgagt cgcgtagaa tagcgaaaca cgggatcggg cgagttcgac   1200
73 cttccgtcgg tctcgccctt tctcgtgttt atacgtttga gcgcactctg agaggttgtc  1260
75 atggcggccg actacga                                               1277
```

RAW SEQUENCE LISTING   DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A   TIME: 14:08:31

Input Set : A:\066001650XPCT.txt
Output Set: N:\CRF4\11222004\I462480A.raw

```
 78 <210> SEQ ID NO: 2
 79 <211> LENGTH: 524
 80 <212> TYPE: DNA
 81 <213> ORGANISM: Mycobacterium tuberculosis
 83 <400> SEQUENCE: 2
 84 ctgcagcagg tgacgtcgtt gttcagccag gtgggcggca ccggcggcgg caacccagcc      60
 86 gacgaggaag ccgcgcagat gggcctgctc ggcaccagtc cgctgtcgaa ccatccgctg     120
 88 gctggtggat caggccccag cgcgggcgcg ggcctgctgc gcgcggagtc gctacctggc     180
 90 gcaggtgggt cgttgacccg cacgccgctg atgtctcagc tgatcgaaaa gccggttgcc     240
 92 ccctcggtga tgccggcggc tgttgccgga tcgtcggtga cgggtggcgc cgctccggtg     300
 94 ggtccgggag cgatgggcca gggttcgcaa tccggcggct ccaccagccc gggtctggtc     360
 96 gcgccggcac cgctcgcgca ggagcgtgaa gaagacgacg aggacgactg ggacgaagag    ·420
 98 gacgactggt gagctcccgt aatgacaaca gacttcccgg ccacccgggc cggaagactt     480
100 gccaacattt tggcgaggaa ggtaaagaga gaaagtagtc cagc                      524
103 <210> SEQ ID NO: 3
104 <211> LENGTH: 481
105 <212> TYPE: DNA
106 <213> ORGANISM: Mycobacterium tuberculosis
108 <400> SEQUENCE: 3
109 ctgcagcagg tgacgtcgtt gttcagccag gtgggcggca ccggcggcgg caacccagcc      60
111 gacgaggaag ccgcgcagat gggcctgctc ggcaccagtc cgctgtcgaa ccatccgctg     120
113 gctggtggat caggccccag cgcgggcgcg ggcctgctgc gcgcggagtc gctacctggc     180
115 gcaggtgggt cgttgacccg cacgccgctg atgtctcagc tgatcgaaaa gccggttgcc     240
117 ccctcggtga tgccggcggc tgttgccgga tcgtcggtga cgggtggcgc cgctccggtg     300
119 ggtccgggag cgatgggcca gggttcgcaa tccggcggct ccaccagccc gggtctggtc     360
121 gcgccggcac cgctcgcgca ggagcgtgaa gaagacgacg aggacgactg ggacgaagag     420
123 gacgactggt gagctcccgt aatgacaaca gacttcccgg ccacccgggc cggaagactt     480
125 g                                                                     481
128 <210> SEQ ID NO: 4
129 <211> LENGTH: 302
130 <212> TYPE: DNA
131 <213> ORGANISM: Mycobacterium tuberculosis
133 <400> SEQUENCE: 4
134 atggcagaga tgaagaccga tgccgctacc ctcgggcagg aggcaggtaa tttcgagcgg      60
136 atctccggcg acctgaaaac ccagatcgac caggtggagt cgacggcagg ttcgttgcag     120
138 ggccagtggc gcggcgcggc ggggacggcc gcccaggccg cggtggtgcg cttccaagaa     180
140 gcagccaata agcagaagca ggaactcgac gagatctcga cgaatattcg tcaggccggc     240
142 gtccaatact cgagggccga cgaggagcag cagcaggcgc tgtcctcgca aatgggcttc     300
144 tg                                                                    302
147 <210> SEQ ID NO: 5
148 <211> LENGTH: 100
149 <212> TYPE: PRT
150 <213> ORGANISM: Mycobacterium tuberculosis
152 <400> SEQUENCE: 5
154 Met Ala Glu Met Lys Thr Asp Ala Ala Thr Leu Gly Gln Glu Ala Gly
155  1               5                  10                  15
158 Asn Phe Glu Arg Ile Ser Gly Asp Leu Lys Thr Gln Ile Asp Gln Val
159                 20                  25                  30
162 Glu Ser Thr Ala Gly Ser Leu Gln Gly Gln Trp Arg Gly Ala Ala Gly
```

RAW SEQUENCE LISTING   DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A   TIME: 14:08:31

Input Set : A:\066001650XPCT.txt
Output Set: N:\CRF4\11222004\I462480A.raw

```
163           35                    40                    45
166 Thr Ala Ala Gln Ala Ala Val Val Arg Phe Gln Glu Ala Ala Asn Lys
167      50                    55                    60
170 Gln Lys Gln Glu Leu Asp Glu Ile Ser Thr Asn Ile Arg Gln Ala Gly
171 65                    70                    75                    80
174 Val Gln Tyr Ser Arg Ala Asp Glu Glu Gln Gln Gln Ala Leu Ser Ser
175                    85                    90                    95
178 Gln Met Gly Phe
179                   100
182 <210> SEQ ID NO: 6
183 <211> LENGTH: 49
184 <212> TYPE: PRT
185 <213> ORGANISM: Mycobacterium tuberculosis
187 <400> SEQUENCE: 6
189 Met Ala Glu Met Lys Thr Asp Ala Ala Thr Leu Gly Gln Glu Ala Gly
190 1              5                    10                    15
193 Asn Phe Glu Arg Ile Ser Gly Asp Leu Lys Thr Gln Ile Asp Gln Val
194                   20                    25                    30
197 Glu Ser Thr Ala Gly Ser Leu Gln Gly Gln Trp Arg Gly Ala Ala Gly
198           35                    40                    45
201 Thr
205 <210> SEQ ID NO: 7
206 <211> LENGTH: 42
207 <212> TYPE: PRT
208 <213> ORGANISM: Mycobacterium tuberculosis
210 <400> SEQUENCE: 7
212 Gln Glu Ala Ala Asn Lys Gln Lys Gln Glu Leu Asp Gly Ile Ser Thr
213 1              5                    10                    15
216 Asn Ile Arg Gln Ala Gly Val Gln Tyr Ser Arg Ala Asp Glu Glu Gln
217                   20                    25                    30
220 Gln Gln Ala Leu Ser Ser Gln Met Gly Phe
221           35                    40
224 <210> SEQ ID NO: 8
225 <211> LENGTH: 21
226 <212> TYPE: PRT
227 <213> ORGANISM: Mycobacterium tuberculosis
229 <400> SEQUENCE: 8
231 Gln Glu Ala Gly Asn Phe Glu Arg Ile Ser Gly Asp Leu Lys Tyr Thr
232 1              5                    10                    15
235 Gln Ile Asp Gln Val
236                   20
239 <210> SEQ ID NO: 9
240 <211> LENGTH: 16
241 <212> TYPE: PRT
242 <213> ORGANISM: Mycobacterium tuberculosis
244 <400> SEQUENCE: 9
246 Gly Asp Leu Lys Thr Gln Ile Asp Gln Val Glu Ser Thr Ala Gly Ser
247 1              5                    10                    15
250 <210> SEQ ID NO: 10
```

RAW SEQUENCE LISTING   DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A   TIME: 14:08:31

        Input Set : A:\066001650XPCT.txt
        Output Set: N:\CRF4\11222004\I462480A.raw

251 <211> LENGTH: 16
252 <212> TYPE: PRT
253 <213> ORGANISM: Mycobacterium tuberculosis
255 <400> SEQUENCE: 10
257 Gly Ser Leu Gln Gly Gln Trp Arg Gly Ala Ala Gly Thr Ala Ala Gln
258 1   5   10   15
261 <210> SEQ ID NO: 11
262 <211> LENGTH: 16
263 <212> TYPE: PRT
264 <213> ORGANISM: Mycobacterium tuberculosis
266 <400> SEQUENCE: 11
268 Gln Glu Ala Ala Asn Lys Gln Lys Gln Glu Leu Asp Glu Ile Ser Thr
269 1   5   10   15
272 <210> SEQ ID NO: 12
273 <211> LENGTH: 28
274 <212> TYPE: PRT
275 <213> ORGANISM: Mycobacterium tuberculosis
277 <400> SEQUENCE: 12
279 Ser Thr Asn Ile Arg Gln Ala Gly Val Gln Tyr Ser Arg Ala Asp Glu
280 1   5   10   15
283 Glu Gln Gln Gln Ala Leu Ser Ser Gln Met Gly Phe
284   20   25
287 <210> SEQ ID NO: 13
288 <211> LENGTH: 16
289 <212> TYPE: PRT
290 <213> ORGANISM: Mycobacterium tuberculosis
292 <400> SEQUENCE: 13
294 Arg Ala Asp Glu Glu Gln Gln Gln Ala Leu Ser Ser Gln Met Gly Phe
295 1   5   10   15
298 <210> SEQ ID NO: 14
299 <211> LENGTH: 21
300 <212> TYPE: DNA
C--> 301 <213> ORGANISM: Artificial/Unknown
304 <220> FEATURE:
305 <221> NAME/KEY: misc_feature
306 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
308 <400> SEQUENCE: 14
309 ctgcagcagg tgacgtcgtt g   21
312 <210> SEQ ID NO: 15
313 <211> LENGTH: 23
314 <212> TYPE: DNA
C--> 315 <213> ORGANISM: Artificial/Unknown
318 <220> FEATURE:
319 <221> NAME/KEY: misc_feature
320 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
322 <400> SEQUENCE: 15
323 ccgggtggcc gggaagtctg tgt   23
326 <210> SEQ ID NO: 16
327 <211> LENGTH: 23

[handwritten: invalid - see item 10 on Error Summary Sheet]

[handwritten: same error]

RAW SEQUENCE LISTING   DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A   TIME: 14:08:31

Input Set : A:\066001650XPCT.txt
Output Set: N:\CRF4\11222004\I462480A.raw

```
        328 <212> TYPE: DNA
C--> 329 <213> ORGANISM: Artificial/Unknown
        332 <220> FEATURE:
        333 <221> NAME/KEY: misc_feature
        334 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
        336 <400> SEQUENCE: 16
        337 actactttct ctttctacct tcc                                           23
        340 <210> SEQ ID NO: 17
        341 <211> LENGTH: 39
        342 <212> TYPE: DNA
C--> 343 <213> ORGANISM: Artificial/Unknown
        346 <220> FEATURE:
        347 <221> NAME/KEY: misc_feature
        348 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
        350 <400> SEQUENCE: 17
        351 gggggatcc ggtaccaggt gacgtcgttg ttcagccag                           39
        354 <210> SEQ ID NO: 18
        355 <211> LENGTH: 39
        356 <212> TYPE: DNA
C--> 357 <213> ORGANISM: Artificial/Unknown
        360 <220> FEATURE:
        361 <221> NAME/KEY: misc_feature
        362 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
        364 <400> SEQUENCE: 18
        365 gggggtacc ggatcctcgt agtcggccgc catgacaac                            39
        368 <210> SEQ ID NO: 19
        369 <211> LENGTH: 31
        370 <212> TYPE: DNA
C--> 371 <213> ORGANISM: Artificial/Unknown
        374 <220> FEATURE:
        375 <221> NAME/KEY: misc_feature
        376 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
        378 <400> SEQUENCE: 19
        379 gggggatcc caggtgacgt cgttgttcag c                                    31
        382 <210> SEQ ID NO: 20
        383 <211> LENGTH: 31
        384 <212> TYPE: DNA
C--> 385 <213> ORGANISM: Artificial/Unknown
        388 <220> FEATURE:
        389 <221> NAME/KEY: misc_feature
        390 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
        392 <400> SEQUENCE: 20
        393 gggggtacc acggtgacgt cgttgttcag c                                    31
        396 <210> SEQ ID NO: 21
        397 <211> LENGTH: 32
        398 <212> TYPE: DNA
C--> 399 <213> ORGANISM: Artificial/Unknown
        402 <220> FEATURE:
        403 <221> NAME/KEY: misc_feature
```

Please correct this in subsequent sequences

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A   TIME: 14:08:32

Input Set : **A:\066001650XPCT.txt**
Output Set: **N:\CRF4\11222004\I462480A.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:34; Xaa Pos. 11

VERIFICATION SUMMARY   DATE: 11/23/2004
PATENT APPLICATION: **US/09/462,480A**   TIME: 14:08:32

Input Set : **A:\066001650XPCT.txt**
Output Set: **N:\CRF4\11222004\I462480A.raw**

```
L:301 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:315 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:329 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:343 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:357 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:18
L:371 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19
L:385 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:20
L:399 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:21
L:413 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:22
L:427 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:23
L:441 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24
L:455 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25
L:469 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26
L:483 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:27
L:532 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29
L:548 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:30
L:564 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:31
L:578 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:32
L:592 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:33
L:620 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
```